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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Minimum DB seq length: 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Maximum bb seq length: Հնննննննն
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N_Geneseq_101002:*
1: /SibS2/j-gdata/
                                                                                                                                                             3: /SIDS2/gogdata/geneseg/genesegn-embt/AN1982 DAT *

(SIDS2/gogdata/geneseg/genesegn-embt/AN1982 DAT *

(SIDS2/gogdata/geneseg/genesegn-embt/AN1992 DAT *

(SI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /Stus2 /gradata/geneseq/gereseqnrenbl/NA1986 (A174
/Stus2 /gradata/geneseq/gereseqnrenbl/NA1981 (A174
                                          /SIDS2/golddia/qeneseq/geneseqn/embl/NA3660.DAT:*
/SIDs2/golddia/qeneseq/geneseqn/embl/NA2e01A.DAT:*
/SIDs2/golddia/qeneseq/geneseqn/embl/NA2e018 EAT.*
/SIDS2/jrglata/grasssl/jrassejn_smbl/NA2002.DAT:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (without alignments)
11055.668 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution

SUMMARIES

Result	1t	Score	Query Match	Query Match Length DB	ព្	ID d1	Description
;	- :	5983	100.0	100 I 200 I To I 20 I	ا د دو د حکم	ARC57) 26	Carrie arowth horm
	2	484	16.9	540	2	ABL57127	Cattle growth horm
a	ند:	434	15.1	7172	i)	AAK84541	Human immune∕hacma
	4	433.2	15.1	(J)	IJ	ABI.57128	Cattle growth horm
a	IJ,	432.4	15.1	£102	FJ FJ	AAK84540	Human immune/haema
	Э,	430.6	15.0	5065	21	AAC69111	Human secreted pro
	7	430.6	15 0	2062	22	AAF33219	Human secreted pro
C	œ	429.6	15.0	4709	2	AAC69112	Human secreted pro
c	9	429.4	15.0	5186	t :	ААНЭ8265	Human EST derived

DNA encoding novel	AA566109	۱.	1.87	14. H	4.4.4	4	٠.
iiപ്പെ immune,'haema	AAK / 36.21	۲. ۲.	34000	14	4.4	44	
Bu⊤an immune,thaema	Cruisday	E	446.14	- 1	4.4	4.	
Genomic sequence #	ABK4 TOR 7	ر د	2503	14 R	424 6	۲۰	3
`	AAK 7 15.72	:	*44.	- ~	4.4	<u>.</u>	
	AAS79076	1,	6475	<u>-</u>	424 5	4	7
	AAS74982	23	5976	14.8	424.6	9	c
PNA en∽odina novel	AA<74900	22	5075	1. 20	24	38	ဂ
DNA encoding novel	AAS74241	23	5976	14.8	424.6	37	c
	AAS73823	23	5976		24	36	ဂ
	AAS72815	Ŋ	5976	14.8	2	G G	c
	AAS70240		5976		24	4	a
ENA cacoding movel	AAS69611		5976		발	ؽ	٠;
lluman testicular a	ABL97611		5446		Ç1	E)	o
	AAE94794		5446		424 6	31	c
Hamai, indiaire, haema	MK67004		5303		2	7	1.1
DNA encoding novel	AAS72315		1936		425	5	C
Genomia nucleotide	AAF54868	13	81145		42E.2	t.,	c
Nucleotide sequenc	AAF54867	22	50000		426 2	27	c
	MAK68332	t)	42233		426.2	10	
Human testicular a	ABL98786	E)	98333		416.1	5	
	AAL07238	() ()	551118		426.2	۲,	
Human testicular a	ABI.98787	ڊء س	22237		426.2	111	
Human teptoductive	AAI.07239	13	11137		420.2	t.	
Homan immone/Kacma	AAK 25.788	1)	3401	14.9	426.2	!	
Human immun∈,'haema	MY75787	(1 (1	3401		426.2	•) >	
encoding	AAS74998	<u>د</u> رو درو	(1.0.6		427.0	۲-	O
	MS73838	ti W	3300	14.9	427.6	18	C
cacoding	AAE72636	5	33C0		427.6		(.
encoding	AAS71653		3300		427.6	16	G
DNA encoding novel	AAS70257		3300	•	27	5	c
DNA encoding novel	MS68177		3300		13	14	a
Buman nenregulin-	MX96733	E)	3068091	4.9	1.7	13	
‼∴man neurequlin-	VVKac540	ر. ر.	15 14 15	<u>.</u>	4.7 7	.	
натап immune/haema	AAK 79598	N	95103	14.9	427 B	1	;
DNA encoding novel	AAS66296	N	6094	14.9	V	10	a

ALI SUMENTS

Pattle growth hermone receptor gene promoter and exon IA region

ps-AUG-21002 (first entry)

ABL57126;

RESULT 1 AB: 57:25

AHL57125 standard; DNA; 2869 BP

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XX

KW Cattle, beel, breeding, growth hormone; somatotropin; receptor; XX

OS Bos taurus.

EH Key Location/qualitiers
ET primer_bind complement (2580..2605)
ET satellite /*tag- a
FT satellite /*tag- b
FT primer_bind 2607.2646
ET primer_bind 2655..2580
ET cach /*tag- c
ET exch /*tag- d
Tybl..286
ET cach /*tag- d
CAC312269-A1.

XX

CAC312269-A1.

XX

LOCATION/QUALITICAL PRODUCT 2312269.
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Matches 2869;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is the promoter and exon 1A region of the bovine growth hormone receptor gene. A polymorphic TG-repeat microsatellite located 90 bp upstream from a major transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              schemes for improvement of cattle performance. Marker assisted selection with the genetic markers avoids the costly theory pictesting associated with traditional breeding schemes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        having fewer than 12 copies of the TG dinucleotide repeat. Use this marker and other genetic markers in linkage disequilibrium with the locus allows implementation of selection and breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           obtaining head of beet cattle with genetic predisposition for altered curross weight, by assaying genetic material from head for polymorphism linked to promoter Pl of expn lA of bovine growth hormone receptor gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2869 HP; 657 A; 640 C; 582 G; 990 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3: Page 41 43; Slpp; English.
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                                               481 GITAGCAGGGICTATAGGTAACTTTTTAAGGAACCTCCTTACTTTTTTTCCATAGTGATT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 CTCGAGGATCCTTGTTGGTGGCCATTTTAAATATAGAAGTGTGTTCATGTCCATGTCCCCAA 60
                                                                                  GITAGGASGACTATAGGAACTTTTTTAAGGAACCTCCTTACTTTTTCCATAGTGATT 540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTCGAGGATCCTTGTTCGTGTCCATTTTAAATATAGAAGTGTGTTCATGTCCATCCCCAA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTGTGAGTCTGTTTTGTAAGTTAAGTTCATTTGTATCATTTTCTTTTTTAGTTTCCACATATA
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                                                                                                                                                                                                                     CALTIGGGG EGCATGTAFCCCFTTCAGTACTAGTTTTTTTCTCTGATATAGACCCCAAGAGTGA
                                                                                                                                                                                                                                                                                               CATTI GGGGTGCATGTATCCTTTECAGTACTAGTTTTTTCTCTGATATATAGCCCAAGAGTGA
                                                                                                                                                                                                                                                                                                                                                                              CTGTCAA1GGACALICAG11AC111CAGG1C11GGC1G11GTAAACAATACTGTAATGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOTAGGTCATCCGTGTTGCTGCAGATGACATTATTTCATTCTTTTTAATGGCCGAGTAAT
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                                                                                                                                                                                                                                                        Similarity
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100.0%, Fred. M., 0
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                            AGGTTTATATAGGAAAGCAAGCAAGACCAAGAATCTACTGCCAAGCGGTGACCAAGAAACG
                                                                                                                                                                                                     GCTGGCCCCATCCTGATGATGATCCTTCTGATCCTTCCAGCCCAGCCCTTCGCCTTTCACTTCACT
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The present sequence is the promotor and exch IA region of the bovine growth hormone receptor gene. A polymorphic TG repeat microsatellite located 90 bp upstream from a major transcription start site in the gene is associated with average weaning weight

Example 2; Fig

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Best Local Similarity 99.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         schemes for improvement of cattle performance. Other genetic markers may include polymorphisms such as the GTA polymorphism site in exon IA. The A allele (lound in indicine cattle) contains a Dral restriction site that is not present in the G allele (lound in taurine cattle). This difference can be used in a bookkybtle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          assay to distinguish the respective alleles. The 2 T/C upstrea polymorphic sites could similarly be used. Marker assisted selection with the denetic markers avoids the costly phenotypic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hating tewer than 12 copies of the T3 dimodeolide repeat. One
this marker and other genetic markers in linkage disequilibrium
with the locus allows implementation of selection and breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and carcass weight of cattle. Cattle having at least 12, and preferably 16-20, copies of the TG dinucleotide repeat marker show increased carcass or weaning weight compared with cattle
   Homo saprens
                                                               Ruman immute/harmatopoietic antigen genomic sequence SEC II No. 39353.
                                                                                      07 NOV 2001 (first entry)
                                                                                                                                          AAK84541 Standard; DNA; 7172 BP.
                                                                                                                                                                                                                   2854 AGGAAGTITTAGAGGTC 2869
                                                                                                                                                                                                                                                                                                      2/34 CTACTS-SAAGCGGTGACAAAAAAGTTCACCATATTCCTCCAACCCGCACTGTT 461 CTACTGCCAAGCAAGTGGTCAAGAAAAGTTCACCATATTCCTCCAACCCGCACCGGTT
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                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 540 BP; 124 A; 124 G; 146 G; 148 T; 0 other;
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                           Similaria, pene therapy, carriae, metantanio,
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                                                                                                                                                                                                                                                       instance hachalapoletic, lanuncyhaen n gedetic antigen
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Pred No 3 9c-104;
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Matches

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CC AAK54951 to AAK64702 encode the human immune/haematopoletic antiqen (1) CC antivolation actid sequences given in AAM62170 to AAM61921. (1) have cytostatic CC activity, and can be used in gene therapy and various production. (1) CC proteins and polymoclecides may be used in the propriate (1) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by recutivity of (1) by expressing inactive proteins or to CC sugliament the activity of (1) by expressing inactive proteins or to CC sugliament the patients own production of (1). Additionally, (1) CC pulymoclecides may be used to produce the secreted (1), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (1) proteins and polymoclecides may be used to prevent, aliquose and treat immune/haematupolicide derived cells. AAK64703 concerns and cancer metastasus of haematopolicide antique genomic consert execution the present invention. AAK64704 to AAK67644 represent the present the protein of the AAK64763 and AAM62169.
                       Query Match
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useful for proventing, diagnosing and/or treating cancers and
metastasis
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                                                                                                                  represent sequences used in the exemplification of the present invention
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Halo CS;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2375 ITANATTOGACAACIGIOGGIGIGITCANGGGIGIGAGGGIAIGGGGIGIGGAAAAIIG 2434
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and marmass weight of cattle. Cattle having at least 12, and preferably 16 20, copies of the TG dinucled life repeat marker show increased carcass or weaning weight compared with cattle having fewer than 12 copies of the TG dinucleotide repeat. Use this marker and other genetic markers in linkage disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bovine growth bormome receptor uche. A polymorphic To repeat microsatellite located 90 bp upstream from a major transcription start site in the gene is associated with average weaning weight
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 522 BP; 124 A; 121 C; 136 G; 141 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2674 STAAITSATTTSSSSASSATTIGSSSAAKSTITAIAIAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2494 CAS...CASGCCTGGCCTTCACTTCAGTGGCCAGAAAACAACCACGCCTGGGAAAAAAAGCGGGGG 2553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2854 AGGANGTITAGACGIC 2869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2734 CTACTGCCAAGCGGTGACCAAGAAACGTTCACCATATTCCTCCTCCAACCCCGCACTGTT 279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2554 wiidolffAfoloAdocAAfdedfleffoffeffeftAAfdfffffefofac Addligfofof 261:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 with the locus allows implementation of selection and breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is the promoter and exon 1A region of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Fig 3; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                403
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E 1
                                                                                                                                                                        463 ACCAACTITAAACCTC 478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               343 CTACIGCGAAGGGGTGACCAAGAAAGTICACCATATICCTCCCCCCAACCCCCAACTGTI 402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       y Match 15.1%; Score 433.2; 18 24; Length 522; Local Similarity 95.6%; Prod. No. 3.50-92;
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AAK84540; 07-NOV-2001

(first entry)

AAK84540 standard; DNA; 6102 BP

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 15.1%;
Host Local Similarity 70.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression by rectifying mutations or deletions in a patient's genome that affect the activity of () by expressing inactive proteins or to supplement the patients own production of (). Additionally, () polynomials may be used to produce the secreted (), by inserting the nucleic acids into a host cell and culturing the cell to express the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid sequences given in AAMB2170 to AAM91921. (I) have cytostalic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynocicotides may be used in the prevention, diagnosis and treatment of discases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased
5078 TATETECGACATCCTCCAGCACCTGCTGTTTTCCTGACTTTTTAATGATTGCCATTCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1804 CHITTOTOCACACCCTICTCYAGGATTTATTATTTCTAGACTTTTGGATCGCAGCCAATCT 1863
                                                                                                                                                                                                                                                                                                        5198 CAGTAATGGGATGGCTGGGTGAAATGGTATFTCTAGTTCTAGATCCCTGAGGAATGGCCA 5139
                                                                                                                                                                                                                                                                                                                                                                           1604 CAGCAG LOGGAG LOGGAG ALAMAG CAGTECTA PELL CAGTELL LITTANG GANTETECKA 1743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1570 ATC IGCTGATGGATGTAGGETGCTTGCATGTCCTGGCTATTATAAACAGTGCTGCGGAT 1629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5378 CTPPPTTATOGCTGCAFAGTATTCCATGGTGTATATGTGCCACATTTTCTTCATCCAGTC 5319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1450 FECACTOTIATAATAGGCTOGAGTTGATGGAGCTGATTAGAACTGATTGAAATGTATT 1509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5498 TOTTCAATICCCACCIAGGAGAGAGAATALGGGGGGTTIGGTTHITIGTTCCTGCGATAG 5439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1999 THE TANANT CONTAINATAIN OF TANANTAINATAIL TAIL THE CONTROL OF THE TANANTAIN THE T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to AAK87694 represent human immune/haemalopoletic antiqen genomic sequences from the present invention. AAK54942 to AAK54950 and AAM82169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancers and cancer metastases of baematopoietic derived cells. AAK54703
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein. (1) proteins and polynomicotides may be used to prevent, diadnose and treat immune/haematopoietic-related diseases, especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       = 2 2
                                                                                                                                              5.138 CACTGACTEC ACAMEGO PROTACTAGETTA CAGEGO ACCAACAO FOTAAAAAG TOTTCO
                                                                                                                                                                                                                                                                                                                                                                                                                                                      5258 AAACATACGIGIGGA AIGIGICTITATAGGAGAGAGALITATAG ICCIIIGGGTATATACC 5199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1630 GAACAFFEGGGFRACACGFCFCFFFFCFFGGFFF ----FCCFCAGFGFGFAFGCC 1683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 6102 BP; 2385 A; 1313 C; 1225 C; 1128 T; 1 Sther;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      represent sequences used in the exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure: SEQ ID NO 39352; 3071pp + Sequence Listing; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acids encoding human immune/hematopoietic autigen polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPT, 2001 483426/52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JAN-2001;
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QΥ
        the fusion protein as compared to the human protein only. The genes proteins are useful for preventing, ameliorating or treating medical conditions, e.g. by protein or gene therapy. The genes are isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4838
                                                          The invention relate to the isolation of genes AAC69084-C69119 encoding 27 human secreted proteins AAS37984-B38019. The genes can be used to generate lusion proteins by linking to the gene for the human immunoglobulin G-Fc portion (AAC69075) for increasing the stability of
                                                                                                                                                                              New nucleic acid molecules encoding 27 human secreted proteins for diagnosing, preventing, treating or ameliorating medical conditions and used as food additions or preservatives.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cytastatic, immun.suppressive, montropic, accompanied ive; antiviral; antiallergic; hepatotropic; antidiabetic; antintlammatory; antidicer; vulnerary, anticonvulsant, antibacterial, antitungal, antiparasitic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 - SEP - 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W0200055371-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human secreted protein gene 3 clone HPJCX13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAC69111 standard; DNA: 5065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4778 TCCCAGGTTGTAGGTTGCCTGTLCACTCTGATGGTAGTCTCTTTTGCTGTGCAGAAGCTC 4719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2104 TCCCARTCTCAAGGCTGTCTFTTCACCTTGCTAATAGTTTCCTTTGATGTCCAGAAAGTTT 2163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1984 TETACHTCET PGCCCCATFFFFFTGACTFGGCFCGFFTATFFFTCTGGGAGTTGAGCTGTAGC 2043
                                                                                                                                                Claim 1; Page 383-384; 453pp; English.
                                                                                                                                                                                                                                                     P-PSDH; AAB38011.
                                                                                                                                                                                                                                                                                                    Komatsoulis G,
                                                                                                                                                                                                                                                                                                                      Ruben SM,
                                                                                                                                                                                                                                                                                                                                                    (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                     18-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                     16-MAR-2000; 2000WO-US06783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acciding said disease, infection, homan, secreted protein; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31 JAN 2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4958 GATGAGCALLILLI ICAIGTALLILLI IGGGLGGALAAATG FCFFCITTTGAGAAGGGTCTG 4899
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                                                                                                                                                                                                                                                                                                    Latleur DW,
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Moore PA,
                                                                                                                                                                                                                                                                                                  Shi Y, Bir
Olsen HS,
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a range of human tissues disclosed in the specification. The

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Matches 592:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosts, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer, and other cancers of the advend pland, bowe, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) cardiovascular diserders such as myccardial ischaemias: (d) wound healing: (e) neurological diseases e.g. cerebral anoxia and cpllepsy; and (f) infectious diseases such as viral, bacterial, fungal
                                                                                                                                                                                                                                                                                                                                                                                                             1450 THEACHERGHALAALAGEREEASTTTCATCCACCTCATTAGAACTGATTCAAATGTATT 1509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 5085 BF; YEO A; Y48 C; 1046 G; 2017 T; 74 Other:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and parasitic infections.
2163 TTTAAGGTTAATTAGGTCCCATTTGTTTATTTTGCTTTTATTTCCAATATTCTGGGAGG 2222
                                                                                                                                                               2103 CTCCCATTCTCAAGGCTGTCTTTTCACCTTGCTAATAGTTTCCTTTGATGTGCAGAGGCT
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                                                                                                                                                                                                                                                 920 GAGTTCWTTCTACATTCTCCATATTACCCCTTTCTCACATCACTACH (ICCAAAAATTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           500 AAACATACGTGTGTGTGTCTTTATAGCAGCATGATTTATARTCCTTTGGGTATATACC
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Pred. No. 3.4e-91;
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                                                                                                                          immunosmodulatory; antisclerotic; dermatological; immunosuppressive; antiliflammatory; anti-HIV; immunostimulant, cytostatic, cardiant; vascular; antimicrobial; anti-anglogenic; ophthalmological; neuroprotectant; anticonvulsant; nootropic; antialzheimers; antiparkinsenian; and vulcerary. The polyhucluctidus and polypeptides be used in the prevention, diagnosis and treatment of diseases associa with inappropriate polypeptide expression. Disorders that may be prevented, diagnosed and/or treated by the above methods include immun disorders (e.g. multiple sclerosis, systemic lupus erythematosus and buman immuno-deficiency virus (HIV) infections), hyperproliferative discreters of the properties of the province of the procession of the province o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      coronary urteriosclerosis; angiogenic disorder; diabetic retinopathy; corneal graft neovascularisation; neurological disorder; regeneration; Huntington's chorea, Alzheimer's disease, Parkinson's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1100 T 1100
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disorders (e.g. cancers and Gaucher's disease), cardiovascular (e.g. Scimitar syndrome, Chaga's cardiomyopathy and coronary arterioscherosis), angiogenic disorders (e.g. corneal grait neovascularisation and diabetic retinopathy), neurological disc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human secreted proteins given in AAB64882 to AAB64930. AAB64931 to AAB64991 represent human secreted polypeptide sequences and proteins homologous to them, which are given in the exemplification of the pro-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acids encoding 49 human secreted polypeptides, useful for preventing, diagnosing and/or treating e.g. cancers, Parkinson's disease and diabetic retinopathy -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       multiple sclerosis; systemic lupus erythematosus; HIV; infection; hyperproliferative disorder, cancer; Gaucher's disease; wound healing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ophthalmological; neuroprotectant, nootropic; anticonvulsant; vul
antialzheimers, antiparkinsonian, antimicrobial, immune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF33219 standard; cDNA; 5065
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 dermatological; immunosuppressive; antiinflammatory; anti-HIV; immunostimulant; cytostatic; cardiant; vascular; anti-angiogenic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              and cells the genes are expressed in Examples of activities include:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P-PSDH; AAH64888
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1100 F 1100
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                                                                              2004 T 2004
                                                                                                                                                                  1940 CTHAGHHAAHAGAR CAHTTURAAHHIGGEHHIGHIGGAHGCHHIGGIGI
                                                                                                                                                                                                       1954 GATTGGTGGGAAATGGTAWGTGATAGTGG TTTGATTTGGATTAGTGTGATAATGAGTGA 1920
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          | 1984 ATTTAGTTCTTTCGCCCCATTTTTCGATTGGGTCGTTTATTTTTCTGGAGTTGAGCTCTAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1924 TETT AND ADMITTED AT CONTROL AND CARD OF THE LITTLE AND ANALOG OF 1982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1804 CTTTTCTCCACACCCCCCCCCCCACCATTTATTATTATTGTAGACTTTTTGGATCGCAGCCAATCT 186
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1940 GITTIAATOOTOAMAATACIGGATIGGATATATAHALAGAAGGITGTATGGATTG 1560
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                560 CAGUAAT GGGA EGGGUCAAAT GGUATTTCTAGT TCTAGATCCCTGAGGAATCGCCA 619
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                                                                                                                                                                                                                                                                                                                                                  980 CTCCCATTYTCTAGGIIGCC1641GACICIGA1GGIAGLIICTIT11GC164GCAGAAGCT
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New nucleic acid molecules encoding 27 human secreted proteins for diagnosing, preventing, treating or ameliorating medical conditions and used as tood additives or preservatives

Claum i; Page 385-386; 453pp; English.

the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer, and other cancers of the adrenal quad, bone, bone marrow, breast, qastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allerates, autoimmune haemolytic anaemia, autoinmune throititis, diabetes mellitus, crobn's disease, multiple sclerosis, rheumatoid arthritis and ulceralive generate lusion proteins by linking to the gene for the human immunoglobulin G Fc portion (AAC69075) for increasing the stability of the Laston protein as compared to the human protein only. The genes and proteins are useful for preventing, ameliorating or treating medical conditions, e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and epilopsy; and (f) infectious diseases such as viral, bacterial, fungal and parasitic intections. The invention relate to the isolation of genes AAC69084-C69119 encoding 27 human secreted proteins AAH77884-H38019. The genes can be used to

sequence 4709 BF; 1868 A; 990 C; 836 G; 939 T; 76 other;

Best Local Similarity Matches 591, Conserv Query Match Best Local (4424 TTTRC1GAGAAIGAIGKIIIGUAKYITUATGUATGIGGCIAGAAAGGAUAIGAAGTGATG 4365 1450 TTCACTCTGTATAATAGGCTCCAGTTTCATCCACCTCATTAGAACTGATTCAAATGTATT 1509 L390 IIICIAAAIGGGAIAIAIAIAIGGGJIAsIAIAGIGIAIIIAIGIIIIIGGIIGGGAGTTAC 1449 Conscivative 15.0%; Score 429.6; DB 21; Length 4709; 70.3%; Pred. No. 5.60-91; 9, Mismatches 234, Indels 7; Gaps

RESULT 8

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                                                                                                    Human; sheep; plq; cow; fruit fly; yeast; hamster; macaque; horse; tomato; monkey; doq; sea urchin; expressed sequence taq; EST; diagnostics; forensic test; gene mapping; genetic disorder; biodiversity, gene therapy, nutrition, ss.
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                                                                                                                                                                                                Human EST-derived coding sequence SEQ ID NO: 122.
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                                                                                                                                                                                                                                                                        AAH98265;
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4593 CACTGACTTCCACAATGGTTGAACTAGTTTACAGTCCCACCAACAGTGTAAAAGTGTTCC
                                                                                                                                                                             1744 CACTGTTCTCCATAGTGGCTGTACTAGTTTGCATTCCCACCACCACCAGTGTAAGAGGGTTCC
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17 - Trij. - 2000;
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                                   1864 GACTGGTGTGAAATGGTACCTCATAGTGG-TTTGATTTGCATTTCTCTGATAATGAGTGA 1922
                                                                                                                   1804 CTTTTCTCCACACCCTCTCCAGCATTTATTATTTGTAGACTTTTGGATCGCAGCCAATCT 1863
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                                                                             TATTTCTCCACATCCTCTCCAGCACCTGTTGTTTCCTGACTTTTTAATGATTGCCATTCT 4474
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Zhang J, Werhman T;
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quantitating a polypoptide in tissue, as molecular weight markers and as a tool supplement. (11) and its binding partners are useful in medical imaging of sites expressing (11). (1) and (11) are useful for treating
                                                                                                        tor identifying expressed sense. (3) is useful in secondary, reshaight to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or
                                                                                                                                                                                                             polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polymerhodides are also used in diagnostics as expressed sequence tags
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated polynocheolide and encoded polypeptides, useful in
diagnostics, forensies, gene mapping, identification of mutations
responsible for genetic disorders or other traits and to assess
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID No 2100; 103pp; English.
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Note: The sequence data for this patent did not appear in the printed Note:
5125 CITTAGTITAATTAGATGCCATITGTCAATTTTGGTGGCTTTTGGTGGTGT 5066
                                                                                                                                                                                        5545 CACLGACI (CACAGAIGGI I GAACTAGITI ACAGTCCCACCACAACAACTAAAAGTGTTCC
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                                                                                    2163 FFT AAGGTTAATTTAGGTGGGATTTTGTTTATTTTTTTGCTTTTTATTTTCCAATATTCTTGGGAGG 2222
                                                                                                                                                                                                                                                                                                                                                                                                     5245 GAGTTCATTGTAGATTCTGGATATTAGCCCTTTGT;CAGATGAGTTAGGTTGCAAAAAATTTT
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Best Local Similarity
Matches 597; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              supplement the patients own production of (1). Additionally (1) palynucleotides may be used to produce the secreted (1), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (1) puterins and polynucleotides may be used to prevent, diagnose and treat immune/harmatopoietic derived cells. AAK64703 to AAK8764 represent human immune/harmatopoietic darrived cells. Grownic sequences from the present invention. AAK5492 to AAK54950 and AAM82169 represent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5.348 AAACATAGGTGTGGATGTGTCTTTATAGCAGCATGATTTTATAGTCCTTTTGGGTATATACC 5289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                1450 TTCAVTCTGTATAANTAGGTCCAGTTCATCCACCTCATTAGAACTGATTCAAACGTATT 1509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reasment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the artivity of (I) by expressing inactive proteins or to
                                                                                                                                                                            5408
                                                                                                                                                                                                                                                                                                                                                                1510 CHTTEHAMIGGERGAGTAATACHECAHHGTATATATGTACGACAGCHHGTTATGCATTC 1569
                                                                                                                                                                                                                                                                                                                                                                                                                            5528 TETRACTIGAGAATIGA FIGOTIFFICOAATTIFCATIGCATGTCCCTTACAAAGGATATGAACTICATC 5469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 50196 MP; 14643 A; 10221 C; 10905 C; 14426 C; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                activity, and can be used in gene therapy and vaccine production. (1) proteins and polynucleotides may be used in the prevention, diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 34410; 3071pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acids encoding human immune/homatopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating canadrs and
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F-Ps(R: AAGH7900, AAGH7901, AAGH7902, AAGH7903, AAGH7904, AAG67905, AAG67906, AAG67907, AAG67908, AAG67909, AAG67910, AAG67911, AAG67912, AAG67914, AAG67914, AAG67918, AAG679194, AAG679921, AAG67993, AAG67993, AAG67993, AAG679921, AAG67993, AAG679921, A
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                                                                                                                                 2001-550179/61.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CACTGACTTCCACAATGGTTGAACTAGTTTACAGTCCCACCAACAGTGTAAAAAGTGTTCC 5169
                                                                                                                                                                                                Steinthorsdottir V,
                                                                                                                                                                                                Gulcher JR;
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Dr 1422160
1422780 AACTGCTGTGACATGATATCTCATAGTGCTTTTGATTTTGCATTTCTCTCATCCCCAGTCA 1422839
                                                                                                                                                                                                                                                                     1422660 CACTGACTTCCACAATGGTTGAACTAGTTTACAGTCCCACCAACAGTGTAAAAAGTGTTCC 1422719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1422540 AAACATACGTGTGCATGTCTTTATAGCAGCATGATTTATAGTCCTTTGGGTATATACC 1422599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1422480 TATCATTGTTGGACATTTGGTTTGGTTCCAAGTCTTTGCTATTGTGAATAGTGCCGCAAT 1422539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1422420 CTTTTTWATGGCTGCATACWATTGCATGGTGTATATGTGCCACATWTTCTTAATCCAGTC 1422479
                                                                                                                                      1422729 TATTTCTCCACATCCTCTCCAGCACCTGTTGTTTCCTGACTTTTTAATGATTGCCATTCT 1422779
                                                                                                                                                                                                                                                                                                                                                                                                                1422600 CAGHAATGGGATGGCTGGGTCAAATGGTAFFTCTAGATTCTTAGATCCCTGAGGAATCGCCA 1422659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              patients may be in need of resturative therapy. The NRGIAGI polypeptides may also be used as antigens in the production of antibodies against NRGIAGI and in assays to identify modulators of NRGIAGI expression and activity. Anti-NRGIAGI antibodies and antagonists may also be used to down regulate expression and activity. Anti-NRGIAGI antibodies may
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of NRGIAGI by expressing inactive proteins or to supplement the patients own production of NRGIAGI. Additionally, the gene may be used to produce NRGIAGI polypeptides, by inserting the nucleic acids into a just cell and culturing the cell to express the protein. The gene may also be used as DNA probes and primers in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NRGIAGI) of the invention. The NRGIAGI gene is also referred to as the human Schizophrenia gene. The invention also relates to fragments or variants of the gene and the NRGIAGI polypeptides they encode. The NRGIAGI nucleic acids and polypeptides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate NRGIAGI expression. For example, they may be used to treat disorders associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1684 CAGCAGTGGGGTTGGGATCATAAGGCAGTTTTATTTCCAGTTTTTAAGGAATCTCCA 1743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1390 IIIUIAAATUCUATATATATGUTTAGUTATATGTTTTTUUTTUUTTAU 1449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         down regulate expression and activity. Anti-NPG1AG1 antibodies may also be used as diagnostic agents for detecting the presence of NRG1AG1 polypeptides in samples. NRG1AG1 Is associated with schizophrenia which
                                                                 1864 GACTGGTGTGAAATGGTAC TCATAGTGGCTTGGATTTGCATTTCTGATAATGAGTGA
                                                                                                                                                                                                    1804 CTTTTCTCCACACCCTCTCCACCATTTATTATTTGTACACTTTTGGATCGCAGCCAATCT 1863
                                                                                                                                                                                                                                                                                                                                          1744 CACTGTTCTCCATAGTGGCTGTACTAGTTTGCATTCCCAGCAGCAGTGTAAGAGGGGTTCC 1803
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1630 GAACATTGGGGTACACGTGTCTCTTTCCCCTTCTGGTT-----TCCTCAGTGTGTATCCC 1683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1450 TTCACTCTGTATAATAGGCTCCAGTTTCATCCACCTCATTAGAACTGATTCAAATGTATT 1509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1503400 BP: 452487 A; 281874 C; 288924 G; 480092 T; 1373 offer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        may be prevented, diagnosed and/or treated by the above methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 90-501; 750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        preventing diagnosing and treating schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAG67927. AAG57928, AAG57929, AAG57930, AAG57931, AAG57932, AAG57933, AAG67934, AAG67935, AAG67936, AAG67937.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence represents the human neuregulin-1 associated gene 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATCTGCTGATGGACATCTAGGTTGCTTCCATGTCCTGGCTATTATAAACAGTGCTGCGAT 1629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TGTTCAATTCCCACCTATGAGTGAAAATATGCGGTGTTTGGGTTTTTTTGCGATAG 1422359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  associated gene 1 nucleic acids and fragments, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14.98;
71.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 427.8; DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. 1.3e-89
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Bb 1422960 GAGTICALLGLAGALICUGGALALFAGCCCFILGLCAGALGACTAGGTTGCGAAAATTTT 1423019
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XPXXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EXEXT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P-PSDB; AAG67938, AAG67939, AAG67940, AAG67941, AAG67942, AAG67944, AAG67945, AAG67946, AAG679947, AAG67948, AAG67954, AAG67954, AAG67955, AAG67955, AAG67955, AAG67955, AAG67955, AAG67955, AAG67956, AAG67956, AAG67956, AAG67956, AAG67956, AAG67956, AAG67966, AAG67963, AAG67964, AAG67965, AAG67963, AAG67964, AAG67971, AAG67972, AAG67973, AAG67958, AAG67952, AAG67963, AAG67970, AAG67971, AAG67972, AAG67973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17 DEC-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAK96733;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK96733 standard; DNA; 1503900 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2043 GAGTIGETIGIATALITIFIGAGAITAGTIGTTIGTEGGTIGETTECATITGETATTATITT 2102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1983 ATHAGHICHHIGGCGCAHITHTTGATTGGGTCGTTTATTTTTTTGGGAGTGGAGCTGTAC
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                                                                                                                                                   Disclosure; Page 345-756; 756pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                         Stetansson H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (DECO-) DECODE GENETICS EHF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-FEB-2000; 2000US-0515716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W0200164877-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; neuregulin 1 gene, schizophrenia, gene therapy; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human neuregulin-1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      z8 EEB zuul, j881Wc USU6377
                                                                                                                                                                                                          Neuregulin 1 nucleic acids and proteins useful for
                                                                                                                                                                                                                                                 AAG67974, AAG67975
                                                                                                                                                                                                                                                                                                                                                                                     2001-514841/56
                                                                                                                                                                                        and treating schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                       Steinthorsdottir V, Gulcher JR;
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and treatment of diseases associated with inappropriate neurequiin 1 expression, such as schizophrenia. For example they may be used to treat disorders associated with decreased neurequiin 1 expression by rectifying

The invention also relates to fragments or variants of the neurogulin lighter. The gene and its proteins may be used in the prevention, diagnosis

This sequence represents the human neuregulin i gene of the invention

mutations or deletions in a patient's genome that affect the activity of

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DE 1422900 GERARGECTICACCACTITITION GOOGLIGHT HITTECTIGIAAATTTGTET 1422959
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                                                                                                                                    14.22960 GAGUTCATTGTAGATTGTGGATATTAGCCCCTTTGTCAGATGAGTAGGTTGGGAAAATTTT 14.23019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14.22600 CASTAATGGGATGGGTGAAATGGTATTTCTAGTTCTAGATGCCTGAGGAATCGCCA 1422659
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2163 FITTAACCTTTAATTTAGGTCCCATTTTGTTTTATTTTTTTGCTTTTATTTTCCAATTATTCTGGGACG 2222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       204 CACTESTICITATATITITISASATIASTISTITOTESSTISETTENISTISETATIATITT 1102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1983 AC PEACHPROSPERATORATION PROPERTY AT PROCESS AND A STATE OF A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1864 GACTGGTGTGAAATGGTAGGTGATAGEGG-TFFGAFFFGCAFFTCTCTGAATAATGAGTGA 1922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1804 CHIPTCHCCACACCCHCHCCAGCAFFFAFTAFFTAFFTGFACACTFFFGGAFCGCAGCCAAFCF 1863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1630 CAACATTGGGGTACACCTCTCTCTTCCCTTTCTGGTT-----TGGTGAGTGTGTATGGC 1683
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           presence of neurogulin 1 in samples.
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Local Similarity 71.0%;
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                                                                                                                                                                                            quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imading of sites expressing (II). (I) and (II) are useful to treating disorders involving aberrant protein expression or biological activity. The golypeptide and polynuclectide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations responsible for generic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated polynucleotide and encoded polypeptides, useful in diagnostics, torensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess the content of the con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCK) primers, oligomers, and for chromosome polymer mapping, and in recombinant production of (II). The and gene mapping, and in recombinant production of (II).
                                                                                                                       amino acid sequences. AAS64197-AAS94564 represent novel human diagnostic coding sequences of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAR-2000; 2000US-0540217
23-AUC 2000; 2000US 0649167
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                                          specification, but was obtained in electronic format directly from Wipo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to restore normal activity of (II) or to treat disease states involving (II) (II) is useful for generating antibodies against it, detecting or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polymocheolides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy technique
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to isolated polymucleotide (I) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1: SEQ ID No 3981; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P-PSDH; AHC03990.
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ftp.wipo.int/pub/published_pct_sequences.
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                                                                                 The sequence data for this patent did not appear in the printed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAS70257 standard; cDNA; 3300
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                                                                                                                                                Human; chromosome mapping, gene mapping; gene therapy; forensic;
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                                                                                                        food supplement, medical imaging, diagnostic, genetic disorder, ss
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The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oliquents, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynuclectide sequences have applications in disorders remained acceptances there applications of mutations.
                                                         diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and amino acid sequences. AAS64197.AAS94564 represent movel human diagnostic coding sequences of the Invention.

Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID No 6061; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated polynucleotide and encoded polynoptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other trails and to assess
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23-AMG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-MAR-2001;
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                                       at ftp.wipo.int/pub/published_pct_sequences.
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Query Match Best Local Similarity Matches 595; conserv Conservative 14.9%, Store 427.6, SB 2 71.2%; Pred. No. 1.5e-90; 0: Mismatches 234, Indels SB 23, Length 3300: Saps

Sequence 3300 BP; 1355 A; 711 C; 566 G; 668 T; 0 other;

Ϋ́ QY Ş g d Ŋ 밁 Ë 1395 AAATCCCATATATATGCGTTAGTATACTGTTTTATGTTTTTCCTTCTGGCTTACTTCAC 1454 1689 GTGGGGTTGCTGGATCATAAGGCAGTTCTATTTCCAGTTTTTTAAGGAATCTCCACACTG 1748 3178 THATGGCTGCATAGTATTCCATGGTGTATATGTGCGAGATTTTCTTANTCCAGTCTATCA 3119 1455 TOTGTATAATAGGOTGOAGTTTCATCOAGCTGATTAGAAGTGATTCAAATGTATTCTTTT 1514 3058 HACGEGEGEGEGETTTATAGCAGCATGATTTATAGTCCTTTEGGGEATATAGCCCAGTA 2999 3118 TIGTIGGACATITGGGTTGGTTCCAAGTCTTTGCTATTGTGAATAATGCCGCAATAAACA 3059 1515 TAATGGCTGAGTAATACTCCATTGTGTATATGTACCACAGCTTTCTTATCCATTCATCTG 1574 TGAGAATGATGTTCCAATTTCATCCATGTCCCTACAAAGGACATGAACTCATTTT 3179 TTGGGGTACACGTGTCTCTTTCCCTTCTGGTT-----TCCTCAGTGTGTGTATGCCCAGCA 1688 CTGATGGACATCTAGGTTGCTTCCATGTCCTGGCTATTATAAACAGTGCTGCGATGAACA 1634 ANTICC/ACCIATERSTERGAMIAISCEGIGITITGGITTITTGITCITGISATASTITAC 3239

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                                  2578 ATTITICITAGOTTICOCTICTICACTICTICATICGTAGTTTCTTCTTCTCTCTGCAGAAGCTCTTTA 2519
                                                   2108 ATTCTCAACGCTCTTTTTCACCTTGCTAATAGTTTCCTTTTGATGTGCAGAAGCTTTTAA 2167
                                                                                                                                                                                                               2878 CHCCACATCCTCTCCACCACCTCTTCTTTCCTGACTTTTTAATGATTGCCATTCTAACTC 2819
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Search completed: January 14, 2003, 14:17:41 Job time : 9188.41 secs